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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
08/936,657	09/24/1997	FRITZ ECKSTEIN	228213	9976
20306	7590	06/08/2004	EXAMINER	
MCDONNELL BOEHNEN HULBERT & BERGHOFF LLP			SCHULTZ, JAMES	
300 S. WACKER DRIVE			ART UNIT	
32ND FLOOR			PAPER NUMBER	
CHICAGO, IL 60606			1635	

DATE MAILED: 06/08/2004

Please find below and/or attached an Office communication concerning this application or proceeding.



UNITED STATES DEPARTMENT OF COMMERCE

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APPLICATION NO./ CONTROL NO.	FILING DATE	FIRST NAMED INVENTOR / PATENT IN REEXAMINATION	ATTORNEY DOCKET NO.
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EXAMINER

ART UNIT	PAPER
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20040604

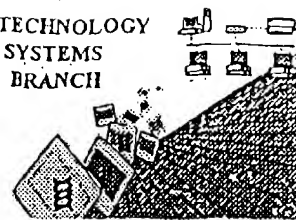
DATE MAILED:

Please find below and/or attached an Office communication concerning this application or proceeding.

Commissioner for Patents

Attached please find an error report generated while attempting to process the computer readable format (CRF) sequence listing submitted March 22, 2004. Sequences 12 and 13 define "N" as being of variable length, which is improper, since variable length entries are not permitted. Correction is required.

JOHN L. LeGUYADER
SUPERVISORY PATENT EXAMINER
TECHNOLOGY CENTER 1600



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 08/936,657A
Source: IFW/6
Date Processed by STIC: 3/29/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/efb/efs/downloads/documents.htm>> , EFS Submission User Manual - cPAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 4B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 08/936,657A
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 _____ Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 _____ Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 _____ Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.	
4 _____ Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 _____ Variable Length	Sequence(s) <u>12-13</u> contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 _____ PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 _____ Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 _____ Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 _____ Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 _____ Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11 _____ Use of <220>	Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12 _____ PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 _____ Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid	



IFW16

RAW SEQUENCE LISTING

PATENT APPLICATION: US/08/936,657A

DATE: 03/29/2004

TIME: 10:30:20

Input Set : A:\00-838-A.SequenceListing.txt

Output Set: N:\CRF4\03292004\H936657A.raw

3 <110> APPLICANT: Sirna Therapeutics, Inc.
 5 <120> TITLE OF INVENTION: Modified Ribozymes
 7 <130> FILE REFERENCE: MBHB 00-838-A; 228/213
 9 <140> CURRENT APPLICATION NUMBER: US 08/936657A
 10 <141> CURRENT FILING DATE: 1997-09-24
 12 <150> PRIOR APPLICATION NUMBER: US 07/965,411
 13 <151> PRIOR FILING DATE: 1993-08-09
 15 <160> NUMBER OF SEQ ID NOS: 15
 17 <170> SOFTWARE: PatentIn version 3.1
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 32
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Artificial Sequence
 24 <220> FEATURE:
 25 <223> OTHER INFORMATION: Ribozyme
 27 <400> SEQUENCE: 1
 28 gatatacctga ctccctatag tgagtcgtat ta 32
 31 <210> SEQ ID NO: 2
 32 <211> LENGTH: 36
 33 <212> TYPE: DNA
 34 <213> ORGANISM: Artificial Sequence
 36 <220> FEATURE:
 37 <223> OTHER INFORMATION: Ribozyme
 39 <400> SEQUENCE: 2
 40 taatacgact cactataggg agtcaggata tctgca 36
 43 <210> SEQ ID NO: 3
 44 <211> LENGTH: 51
 45 <212> TYPE: DNA
 46 <213> ORGANISM: Artificial Sequence
 48 <220> FEATURE:
 49 <223> OTHER INFORMATION: Ribozyme
 51 <400> SEQUENCE: 3
 52 ggagtttcgg cctaacggcc tcatacagagg accctatagt gagtcgtatt a 51
 55 <210> SEQ ID NO: 4
 56 <211> LENGTH: 55
 57 <212> TYPE: DNA
 58 <213> ORGANISM: Artificial Sequence
 60 <220> FEATURE:
 61 <223> OTHER INFORMATION: Ribozyme
 63 <400> SEQUENCE: 4
 64 taatacgact cactataggg tectctgatg aggccgtag gccgaaactc ctgca 55
 67 <210> SEQ ID NO: 5
 68 <211> LENGTH: 34

Does Not Comply
Corrected Diskette Needed

P.3

RAW SEQUENCE LISTING

DATE: 03/29/2004

PATENT APPLICATION: US/08/936,657A

TIME: 10:30:20

Input Set : A:\00-838-A.SequenceListing.txt

Output Set: N:\CRF4\03292004\H936657A.raw

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69 <212> TYPE: DNA
70 <213> ORGANISM: Artificial Sequence
72 <220> FEATURE:
73 <223> OTHER INFORMATION: Ribozyme
75 <400> SEQUENCE: 5
76 ggguccucug augaggccgu uaggccgaaa cucc 34
79 <210> SEQ ID NO: 6
80 <211> LENGTH: 34
81 <212> TYPE: DNA
82 <213> ORGANISM: Artificial Sequence
84 <220> FEATURE:
85 <223> OTHER INFORMATION: Ribozyme
87 <220> FEATURE:
88 <221> NAME/KEY: misc_feature
89 <223> OTHER INFORMATION: All U nucleic acids are 2'-fluoro derivatives.
92 <400> SEQUENCE: 6
93 ggguccucug augaggccgu uaggccgaaa cucc 34
96 <210> SEQ ID NO: 7
97 <211> LENGTH: 34
98 <212> TYPE: DNA
99 <213> ORGANISM: Artificial Sequence
101 <220> FEATURE:
102 <223> OTHER INFORMATION: Ribozyme
104 <220> FEATURE:
105 <221> NAME/KEY: misc_feature
106 <223> OTHER INFORMATION: All U nucleic acids are 2'-amino derivatives.
109 <400> SEQUENCE: 7
110 ggguccucug augaggccgu uaggccgaaa cucc 34
113 <210> SEQ ID NO: 8
114 <211> LENGTH: 12
115 <212> TYPE: RNA
116 <213> ORGANISM: Artificial Sequence
118 <220> FEATURE:
119 <223> OTHER INFORMATION: Substrate
121 <400> SEQUENCE: 8
122 gggagucagg au 12
125 <210> SEQ ID NO: 9
126 <211> LENGTH: 12
127 <212> TYPE: RNA
128 <213> ORGANISM: Artificial Sequence
130 <220> FEATURE:
131 <223> OTHER INFORMATION: Substrate
133 <220> FEATURE:
134 <221> NAME/KEY: misc_feature
135 <223> OTHER INFORMATION: All U nucleotides are 2'-fluoro derivatives
138 <400> SEQUENCE: 9
139 gggagucagg au 12
142 <210> SEQ ID NO: 10
143 <211> LENGTH: 12

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RAW SEQUENCE LISTING

DATE: 03/29/2004

PATENT APPLICATION: US/08/936,657A

TIME: 10:30:20

Input Set : A:\00-838-A.SequenceListing.txt

Output Set: N:\CRF4\03292004\H936657A.raw

144 <212> TYPE: RNA
 145 <213> ORGANISM: Artificial Sequence
 147 <220> FEATURE:
 148 <223> OTHER INFORMATION: Substrate
 150 <220> FEATURE:
 151 <221> NAME/KEY: misc_feature
 152 <222> LOCATION: (7)..(7)
 153 <223> OTHER INFORMATION: 2'-fluoro derivative
 156 <400> SEQUENCE: 10

157 gggagucagg au

12

160 <210> SEQ ID NO: 11

161 <211> LENGTH: 12

162 <212> TYPE: RNA

163 <213> ORGANISM: Artificial Sequence

165 <220> FEATURE:

166 <223> OTHER INFORMATION: Substrate

168 <220> FEATURE:

169 <221> NAME/KEY: misc_feature

170 <222> LOCATION: (6)..(6)

171 <223> OTHER INFORMATION: 2'-amino derivative

174 <400> SEQUENCE: 11

175 gggagucagg au

12

178 <210> SEQ ID NO: 12

179 <211> LENGTH: 13

180 <212> TYPE: DNA

181 <213> ORGANISM: Artificial Sequence

183 <220> FEATURE:

184 <223> OTHER INFORMATION: Substrate

186 <220> FEATURE:

187 <221> NAME/KEY: misc_feature

188 <222> LOCATION: (13)..(13)

189 <223> OTHER INFORMATION: N stands for any one or more A, T, G, C.

192 <400> SEQUENCE: 12

W--> 193 gggagucagg aun

13

196 <210> SEQ ID NO: 13

197 <211> LENGTH: 13

198 <212> TYPE: DNA

199 <213> ORGANISM: Artificial Sequence

201 <220> FEATURE:

202 <223> OTHER INFORMATION: Substrate

204 <220> FEATURE:

205 <221> NAME/KEY: misc_feature

206 <223> OTHER INFORMATION: All U nucleotides are 2'-fluoro derivatives.

209 <220> FEATURE:

210 <221> NAME/KEY: misc_feature

211 <222> LOCATION: (13)..(13)

212 <223> OTHER INFORMATION: N stands for one or more modified or unmodified nucleotides.

215 <400> SEQUENCE: 13

W--> 216 gggagucagg aun

13

*variable length is not permitted.**See
item 5
on Ena
summary
sheet.**same error*

RAW SEQUENCE LISTING

DATE: 03/29/2004

PATENT APPLICATION: US/08/936,657A

TIME: 10:30:20

Input Set : A:\00-838-A.SequenceListing.txt

Output Set: N:\CRF4\03292004\H936657A.raw

219 <210> SEQ ID NO: 14
220 <211> LENGTH: 12
221 <212> TYPE: DNA
222 <213> ORGANISM: Artificial Sequence
224 <220> FEATURE:
225 <223> OTHER INFORMATION: Substrate
227 <220> FEATURE:
228 <221> NAME/KEY: misc feature
229 <223> OTHER INFORMATION: All U nucleotides are 2'-fluoro derivatives.
232 <400> SEQUENCE: 14
233 gggagucagg au 12
236 <210> SEQ ID NO: 15
237 <211> LENGTH: 36
238 <212> TYPE: DNA
239 <213> ORGANISM: Artificial Sequence
241 <220> FEATURE:
242 <223> OTHER INFORMATION: Ribozyme
244 <400> SEQUENCE: 15
245 cacaacacug augaggccgu uaggccgaaa cgggca 36

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 03/29/2004
PATENT APPLICATION: US/08/936,657A TIME: 10:30:21

Input Set : A:\00-838-A.SequenceListing.txt
Output Set: N:\CRF4\03292004\H936657A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:12; N Pos. 13
Seq#:13; N Pos. 13

VERIFICATION SUMMARY

DATE: 03/29/2004

PATENT APPLICATION: US/08/936,657A

TIME: 10:30:21

Input Set : A:\00-838-A.SequenceListing.txt

Output Set: N:\CRF4\03292004\H936657A.raw

L:193 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:216 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0